SEQUENCE LISTING

<110> Pagratis, Nikos Lochrie, Michael Gold, Larry <120> High Affinity TGFBeta Nucleic Acid Ligands and Inhibitors <130> NEX87 <140> 09/363,939 <141> 1999-07-29 <150> 09/046,247 <151> 1998-03-23 <150> 08/458,424 <151> 1995-06-02 <150> 07/714,131 <151> 1991-06-10 <150> 07/931,473 <151> 1992-08-17 <150> 07/964,624 <151> 1992-10-21 <150> 08/117,991 <151> 1993-09-08 <150> 07/536,428 <151> 1990-06-11 <160> 216 <170> PatentIn Ver. 2.0 <210> 1 <211> 71 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Sequence <220> <221> modified base <222> (1)..(71) <223> Feature N at positions 17-56 is A, C, G or T. <400> 1 atcgtcctcc c

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tcgggcgagt cgtctg
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taatacgact cactataggg aggacgatgc ggnnnnnnnn nnnnnnnnn nnnnnnnnn 60
nnnnnnnn nncagacgac tcgcccga
<210> 5
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<223> Feature N at positions 33-72 is A, C, G or T.
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attatgctga gtgatatccc tcctgctacg ccnnnnnnn nnnnnnnnn nnnnnnnnn 60
nnnnnnnn nngtctgctg agcgggct
<210> 6
<211> 71
<212> RNA
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<223> All pyrimidines are 2'F; feature N at positions
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71
gacucgcccg a
<210> 7
<211> 65
<212> DNA
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<223> Feature N at positions 16-49 is SEQ ID NO: 115
     and A, C, G or T.
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gggaggacga tgcggnnnnn nnnnnnnnn nnnnnnnnn nnnnnnnnn agacgactcg 60
cccga
<210> 8
<211> 82
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<223> Feature N at positions 33-66 is SEQ ID NO: 115 and
     A, C, G or T.
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taatacgact cactataggg aggacgatgc ggnnnnnnnn nnnnnnnnn nnnnnnnnn 60
nnnnncaga cgactcgccc ga
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<400> 9
attatgctga gtgatatccc tcctgctacg ccnnnnnnn nnnnnnnnn nnnnnnnnn 60
nnnnnngtct gctgagcggg ct
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      16-49 is SEQ ID NO: 115 and A, 2'-FC, G or 2'-FU.
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gggaggacga ugcggnnnnn nnnnnnnnn nnnnnnnnn nnnnnnnnn agacgacucg 60
<210> 11
<211> 27
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cgcaggatcc taatacgact cactata
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<210> 12
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ggcag	aattc tcatctactt agtcgggcga gtcgtctg		38
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<400> gtgct	14 gcaag gcgattaagt tgg		23
<210><211><212><212><213>	19		
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<400> acttta	15 atgct teeggeteg		19
<210> <211> <212> <213>	19		
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<400>	16		19

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<211> 18
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ggggacaagc ggacttag
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<210> 19
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ccgcaugcuc cuccc
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<210> 22
<211> 71
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gggaggacga ugcggaagua acguuuaagu aaaauucguu cucucgguau uuggccagac 60
gacucgcccg a
<210> 23
<211> 71
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gggaggacga ugcggaagua acguugaagu aaaauucguu cucucggcau uuggccagac 60
gacucgcccg a
<210> 24
<211> 71
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gacucgcccg a
<210> 25
<211> 69
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cucgcccga
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gggaggacga ugcggauaga ucgguccgau aagucuuuca ucuuuaccug gcccccagac 60
gacucgcccg a
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<210> 27
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<223> All pyrimidines are 2'F.
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gggaggacga ugcggaagua acquuqaagu aaaauucquu cucucgguau uuqqccaqac 60
gacucgcccg a
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gggaggacga ugcggacgau ccuuuccuua acauuucauc auuucuccug ugccccagac 60
gacucgcccg g
<210> 29
<211> 71
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gggaggacga ugcgguccau caacaaucuu aucauuaugu uuuuccuucc cgccccagac 60
gacucgcccg a
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<210> 30
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gacucgcccg a
                                                                    71
<210> 31
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gggaggacga ugcgguuccu caauucuucc aucuucauaa uguuucccuu ugccccagac 60
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gacucgcccg a
<210> 32
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gacucgcccg g
<210> 33
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<222> (1)..(71)
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gggaggacga ugcggucuca acgaagaaca ucguuggaua cuguuugucc cgccccagac 60
<210> 34
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gggaggacga ugcgguucag uuuccuucag uuuucguuuc uaauucuugu guccccagac 60
gacucgcccg a
                                                                    71
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gacucgcccg a
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acucgcccga
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gacucgcccg a
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acucgcccga
<210> 43
<211> 71
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gacucgcccg a
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gacucgcccg a
<210> 46
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acucgcccga
<210> 47
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gacucgcccg a
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acucgcccga
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gggaggacga ugcggucuug auccuccuuu gugucuuucu uugucuuccc ugccccagac 60
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acucgcccga
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acucgcccga
<210> 56
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gacucgcccg a
<210> 57
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                                                                    71
gacucgcccg a
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<400> 58
gggaggacga ugcggagaug guacuccauc uccuuuaugu gcccaucgcu guccccagac 60
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71
gacucgcccg a
<210> 59
<211> 61
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                                                                    61
<210> 60
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gacucgcccg a
                                                                    71
<210> 61
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acucgcccga
                                                                    70
<210> 62
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<211> 71
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gacucgcccg a
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gacucgcccg a
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gacucgcccg a
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<212> RNA
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gacucgcccg a
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acucgcccga
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acucgcccga
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acucgcccga
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<210> 90
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<212> RNA
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gguucaggag guuauuacag agucuguaua gcuguacucc cca
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<210> 92
<211> 36
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ggagguuauu acagagucug uauagcugua cucc
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<210> 94
<211> 30
<212> RNA
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ggagguuauu acagagucug uauagcugua cu
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                                                                    32
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<210> 112
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ggagguuauu acagagucug uauagcugua cucccca
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Sequence

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<400> 120
gguuauuaca gagucuguau agcuguacuc c
                                                                    31
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<211> 34
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                                                                    34
<210> 122
<211> 60
<212> PRT
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Ala Leu Asp Thr Asn Tyr Cys Phe Ser Ser Thr Glu Lys Asn Cys Cys
                  5
                                      10
Val Arg Gln Leu Tyr Ile Asp Phe Arg Lys Asp Leu Gly Trp Lys Trp
Ile His Glu Pro Lys Gly Tyr His Ala Asn Phe Cys Leu Gly Pro Cys
                              40
Pro Tyr Ile Trp Ser Leu Asp Thr Gln Tyr Ser Lys
     50
                         55
<210> 123
<211> 60
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 123
Ala Leu Asp Ala Ala Tyr Cys Phe Arg Asn Val Gln Asp Asn Cys Cys
                                      10
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Leu Arg Pro Leu Tyr Ile Asp Phe Lys Arg Asp Leu Gly Trp Lys Trp 20 25 30

Ile His Glu Pro Lys Gly Tyr Asn Ala Asn Phe Cys Ala Gly Ala Cys
35 40 45

Pro Tyr Leu Trp Ser Ser Asp Thr Gln His Ser Arg
50 55 60

<210> 124

<211> 60

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
 Sequence

<400> 124

Ala Leu Asp Thr Asn Tyr Cys Phe Arg Asn Leu Glu Glu Asn Cys Cys 1 5 10 15

Val Arg Pro Leu Tyr Ile Asp Phe Arg Gln Asp Leu Gly Trp Lys Trp 20 25 30

Val His Glu Pro Lys Gly Tyr Tyr Ala Asn Phe Cys Ser Gly Pro Cys
35 40 45

Pro Tyr Leu Arg Ser Ala Asp Thr Thr His Ser Thr 50 55 60

<210> 125

<211> 52

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
 Sequence

<400> 125

Val Leu Ala Leu Tyr Asn Gln His Asn Pro Gly Ala Ser Ala Ala Pro 1 5 10 15

Cys Cys Val Pro Gln Ala Leu Glu Pro Leu Pro Ile Val Tyr Tyr Val 20 25 30

Gly Arg Lys Pro Lys Val Glu Gln Leu Ser Asn Met Ile Val Arg Ser 35 40 45

Cys Lys Cys Ser 50

<210> 126

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                                      10
Cys Cys Val Ser Gln Asp Leu Glu Pro Leu Thr Ile Leu Tyr Tyr Ile
                                  25
Gly Lys Thr Pro Lys Ile Glu Gln Leu Ser Asn Met Ile Val Lys Ser
                              40
Cys Lys Cys Ser
     50
<210> 127
<211> 52
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<400> 127
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Cys Cys Val Pro Gln Asp Leu Glu Pro Leu Thr Ile Leu Tyr Tyr Val
             20
                                  25
                                                      30
Gly Arg Thr Pro Lys Val Glu Gln Leu Ser Asn Met Val Val Lys Ser
         35
                              40
Cys Lys Cys Ser
     50
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<211> 52

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<223> All pyrimidines are 2'F.
<400> 128
ggaaguaacg uuguaguaaa auucguucuc ucggcauuug gccagacgac ucgcccga
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gggaggacga ugcggaagua acguuguagu aaaauucguu cucucggcau uuqqcca
                                                                    57
<210> 130
<211> 44
<212> RNA
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<222> (1)..(44)
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ggaaguaacg uuguaguaaa auucguucuc ucggcauuug gcca
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<210> 131
<211> 38
<212> RNA
<213> Artificial Sequence
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gggaggaugc ggaaguaacg uuguaguaaa auucc
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gggaggaugc ggaaguaacg uuguaguaaa au
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<212> RNA
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<222> (1)..(57)
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                                                                    44
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<211> 44
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<222> (1)..(44)
<223> All pyrimidines are 2'F.
<400> 140
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<210> 141
<211> 29
<212> RNA
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ggggagcggc guuguugaaa aguccgcuu
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<210> 142
<211> 38
<212> RNA
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      Sequence
<220>
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<222> (1)..(33)
<223> All pyrimidines are 2'F.
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<210> 144
<211> 42
<212> RNA
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<223> All pyrimidines are 2'F.
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ggggagcggc guuguuuagu cguauquaua uacuaagucc qc
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<210> 145
<211> 38
<212> RNA
<213> Artificial Sequence
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<222> (1)..(38)
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<210> 146
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<212> RNA
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      Sequence
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<222> (1)..(65)
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cccga
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<222> (1)..(64)
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gggaggacga ugcggaggcg uuauuagaga gucuguauag cucuagcccc agacgacucg 60
ccga
                                                                    64
<210> 148
<211> 65
<212> RNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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<220>
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<210> 149
<211> 65
<212> RNA
<213> Artificial Sequence
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<220>
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<222> (1)..(65)
<223> All pyrimidines are 2'F.
<400> 149
gggaggacga ugcggggagg uuauuauaga gucuguauag cuauaccccc agacgacucg 60
cccga
<210> 150
<211> 65
<212> RNA
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<222> (1)..(65)
<223> All pyrimidines are 2'F.
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cccga
                                                                    65
<210> 151
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<221> modified base
<222> (1)..(65)
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<223> All pyrimidines are 2'F.
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cccga
<210> 152
<211> 65
<212> RNA
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<220>
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<222> (1)..(65)
<223> All pyrimidines are 2'F.
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cccga
<210> 153
<211> 65
<212> RNA
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<220>
<221> modified base
<222> (1)..(65)
<223> All pyrimidines are 2'F.
<400> 153
gggaggacga ugcgggcgga uuaucacaga guauguauag cugugccgcc agacgacucg 60
cccga
                                                                    65
<210> 154
<211> 65
<212> RNA
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<220>
<221> modified base
<222> (1)..(65)
<223> All pyrimidines are 2'F.
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cccga
<210> 155
<211> 65
<212> RNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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<220>
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<222> (1)..(65)
<223> All pyrimidines are 2'F.
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cccga
<210> 156
<211> 65
<212> RNA
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      Sequence
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cccga
<210> 157
<211> 65
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      Sequence
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<222> (1)..(65)
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gggaggacga ugcgggggga cuauuaguga gucuguauag cacuaccccc agacgacucg 60
cccga
                                                                    65
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<210> 158
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<212> RNA
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<222> (1)..(65)
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<400> 158
gggaggacga ugcgggugga uuauuacagc gucuguauau cuguaccccc agacgacucq 60
cccga
                                                                    65
<210> 159
<211> 65
<212> RNA
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<223> All pyrimidines are 2'F.
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cccga
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      Sequence
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<222> (1)..(65)
<223> All pyrimidines are 2'F.
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cccga
<210> 161
<211> 65
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<212> RNA
<213> Artificial Sequence
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      Sequence
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<222> (1)..(65)
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<210> 162
<211> 65
<212> RNA
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<223> All pyrimidines are 2'F.
gggaggacga ugcgggugga uuauuacaga gucuguauag cuguaccccc agacgacucg 60
cccga
                                                                    65
<210> 163
<211> 65
<212> RNA
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<220>
<221> modified base
<222> (1)..(65)
<223> All pyrimidines are 2'F.
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gggaggacga ugcgggggcg uuauuacaga gucuguauag cuguagcccc agacgacucg 60
cccga
                                                                    65
<210> 164
<211> 65
<212> RNA
<213> Artificial Sequence
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      Sequence
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<222> (1)..(65)
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cccga
<210> 165
<211> 65
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<222> (1)..(65)
<223> All pyrimidines are 2'F.
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cccga
<210> 166
<211> 65
<212> RNA
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gggaggacga ugcggggagu uuauuacagc gucuguauau cuguagcccc agacgacucg 60
cccga
                                                                    65
<210> 167
<211> 65 ·
<212> RNA
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Sequence
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<223> All pyrimidines are 2'F.
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cccga
<210> 168
<211> 65
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<222> (1)..(65)
<223> All pyrimidines are 2'F.
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gggaggacga ugcggggugg uuauuagaga gucuguauag cucuacgccc agacgacucg 60
<210> 169
<211> 65
<212> RNA
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      Sequence
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<221> modified base
<222> (1)..(65)
<223> All pyrimidines are 2'F.
gggaggacga ugcgggggga guauuaaaga gucuguauag cuuuaccccc agacgacucg 60
cccga
                                                                    65
<210> 170
<211> 65
<212> RNA
<213> Artificial Sequence
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<220>

Sequence

<220>

<223> Description of Artificial Sequence: Synthetic

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<221> modified base
<222> (1)..(65)
<223> All pyrimidines are 2'F.
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gggaggacga ugcggggagg auauuauaga gucuguauag cuauaccccc agacgacucg 60
cccga
<210> 171
<211> 34
<212> RNA
<213> Artificial Sequence
<220>
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<220>
<221> modified base
<222> (1)..(34)
<223> All pyrimidines are 2'F; a's and g's at positions
      1-5, 8, 11, 13-16, 20, 22, 24-25, 28 and 30 are
      2'-OCH3; linkage at positions 34 and 35 is 3'-3'.
<400> 171
ggagguuauu acagagucug uauaqcuqua cucc
                                                                   34
<210> 172
<211> 34
<212> RNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<220>
<221> modified_base
<222> (1)..(34)
<223> All pyrimidines are 2'F; a's and g's at positions
      11, 13-16, 20, 22, 24-25, 28 and 30 are 2'OCH3;
      linkage at positions 34 and 35 is 3'-3'.
<400> 172
ggagguuauu acagagucug uauagcugua cucc
                                                                   34
<210> 173
<211> 34
<212> RNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Sequence
<220>
<221> modified_base
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<222> (1)..(34)
<223> All pyrimidines are 2'F; a's and g's at positions
      1-4 are 2'OCH3; linkage at positions 34 and 35 is
      3'-3'.
<400> 173
qqaqquuauu acaqaqucuq uauaqcuqua cucc
                                                                   34
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<220>
<221> modified base
<222> (1)..(34)
<223> All pyrimidines are 2'F; a's and g's at positions
      5, 8, and 11 are 2'OCH3; linkage at positions 34
      and 35 is 3'-3'.
<400> 174
ggagguuauu acagagucug uauagcugua cucc
                                                                   34
<210> 175
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<223> Description of Artificial Sequence: Synthetic
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<220>
<221> modified base
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<223> All pyrimidines are 2'F; a' and g's at positions
      13-16 are 2'-OCH3; linkage at positions 34 and 35
      is 3'-3'.
<400> 175
ggagguuauu acagagucug uauagcugua cucc
                                                                   34
<210> 176
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<223> Description of Artificial Sequence: Synthetic
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<220>
<221> modified base
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<222> (1)..(34)
<223> All pyrimidines are 2'F; a's and g's at positions
      20, 22 and 24 are 2'-OCH3; linkage at positions 34
      and 35 is 3'-3'.
<400> 176
ggagguuauu acagagucug uauagcugua cucc
                                                                   34
<210> 177
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<223> Description of Artificial Sequence: Synthetic
      Sequence
<220>
<221> modified base
<222> (1)..(34)
<223> All pyrimidines are 2'F; a's and g's at positions
      25, 28 and 30 are 2'-OCH3; linkage at positions 34
      and 35 is 3'-3'.
<400> 177
ggagguuauu acagagucuq uauaqcuqua cucc
                                                                   34
<210> 178
<211> 34
<212> RNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<220>
<221> modified base
<222> (1)..(34)
<223> All pyrimidines are 2'F; g at position 20 is
      2'-OCH3; linkage at positions 34 and 35 is 3'-3'.
<400> 178
ggagguuauu acagagucug uauagcugua cucc
                                                                   34
<210> 179
<211> 34
<212> RNA
<213> Artificial Sequence
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      Sequence
<220>
<221> modified_base
<222> (1)..(34)
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<223> All pyrimidines are 2'F; a at position 22 is
      2'-OCH3; linkage at positions 34 and 35 is 3'-3'.
<400> 179
ggagguuauu acagagucug uauagcugua cucc
                                                                   34
<210> 180
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<223> Description of Artificial Sequence: Synthetic
      Sequence
<220>
<221> modified base
<222> (1)..(34)
<223> All pyrimidines are 2'F; a at position 24 is
      2'-OCH3; linkage at positions 34 and 35 is 3'-3'.
<400> 180
ggagguuauu acagagucuq uauaqcuqua cucc
                                                                   34
<210> 181
<211> 34
<212> RNA
<213> Artificial Sequence
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      Sequence
<220>
<221> modified_base
<222> (1)..(34)
<223> All pyrimidines are 2'F; a's and g's at positions
      1-5, 8, 11, 25 and 30 are 2'-OCH3; linkage at
      positions 34 and 35 is 3'-3'.
<400> 181
ggagguuauu acagagucug uauagcugua cucc
                                                                   34
<210> 182
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<223> Description of Artificial Sequence: Synthetic
      Sequence
<220>
<221> modified_base
<222> (1)..(34)
<223> All pyrimidines are 2'F; a's and g's at positions
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linkage at positions 34 and 35 is 3'-3'.
<400> 182
ggagguuauu acagagucug uauagcugua cucc
                                                                    34
<210> 183
<211> 34
<212> RNA
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<223> Description of Artificial Sequence: Synthetic
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<220>
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<222> (1)..(34)
<223> All pyrimidines are 2'F; a's and g's at positions
      1-5, 8, 11, 13-16, 20, 24-25, 28 and 30 are
      2'-OCH3; linkage at positions 34 and 35 is 3'-3'.
<400> 183
ggagguuauu acagagucug uauagcugua cucc
                                                                    34
<210> 184
<211> 34
<212> RNA
<213> Artificial Sequence
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      Sequence
<220>
<221> modified base
<222> (1)..(34)
<223> All pyrimidines are 2'F; a's and g's at positions
      1-5, 8, 11, 13-16, 25, 28 and 30 are 2'-OCH3;
      linkage at positions 34 and 35 is 3'-3'.
<400> 184
ggagguuauu acagagucug uauagcugua cucc
                                                                    34
<210> 185
<211> 34
<212> RNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<220>
<221> modified base
<222> (1)..(34)
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1-5, 8, 11, 13-16, 24-25, 28 and 30 are 2'-OCH3;

<223> All pyrimidines are 2'F; a's and g's at positions

```
linkage at positions 34 and 35 is 3'-3'.
<400> 185
ggagguuauu acagagucug uauagcugua cucc
                                                                   34
<210> 186
<211> 34
<212> RNA
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<223> Description of Artificial Sequence: Synthetic
      Sequence
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<221> modified base
<222> (1)..(34)
<223> All pyrimidines are 2'F; a's and g's at positions
      1-4, 13-16, 25, 28 and 30 are 2'-OCH3; linkage at
      positions 34 and 35 is 3'-3'.
<400> 186
ggagguuauu acagagucug uauagcugua cucc
                                                                   34
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<223> Description of Artificial Sequence: Synthetic
      Sequence
<220>
<221> modified base
<222> (1)..(33)
<223> All pyrimidines are 2'F; a's and g's at positions
      1-5, 10, 12-15 24, 27 and 29 are 2'-OCH3; linkage
      at positions 33 and 34 is 3'-3'.
<400> 187
ggagauauua cagagucugu auagcuguac ucc
                                                                   33
<210> 188
<211> 32
<212> RNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
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<220>
<221> modified_base
<222> (1)..(32)
```

1-5, 11, 13-16, 25, 28, and 30 are 2'-OCH3;

<223> All pyrimidines are 2'F; a's and g's at positions

```
<400> 188
gggguuauua cagagucugu auagcuguac cc
                                                                      32
<210> 189
<211> 34
<212> RNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<220>
<221> modified base
<222> (1)..(34)
<223> All pyrimidines are 2'F; linkage at positions 34
      and 35 is 3'-3'.
<400> 189
ugugaauauu agagagucug uauagcucua cccc
                                                                      34
<210> 190
<211> 34
<212> RNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<220>
<221> modified base
<222> (1)..(34)
<223> All pyrimidines are 2'F; a's and g's at positions
      2, 4-5, 8, 11, 13-16, 24-25 \text{ and } 30 \text{ are } 2'-\text{OCH3};
      linkage at positons 34 and 35 is 3'-3'.
<400> 190
ugugaauauu agagagucug uauagcucua cccc
                                                                      34
<210> 191
<211> 34
<212> RNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Sequence
<220>
<221> modified_base
<222> (1)..(34)
```

1-4, 10, 12-15, 24, 27 and 29 are 2'-OCH3; linkage

at positions 32 and 33 is 3'-3'.

```
<223> All pyrimidines are 2'F; a's and g's at positions
      2, 4-6, 8, 11-16, 20, 24-25, and 30 are 2'-OCH3;
      linkage at positions 34 and 35 is 3'-3'.
<400> 191
ugugaauauu agagagucug uauagcucua cccc
                                                                   34
<210> 192
<211> 34
<212> RNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<220>
<221> modified base
<222> (1)..(34)
<223> All pyrimidines are 2'F; a's and g's at positions
      2, 4, 11, 13-16, 25 and 30 are 2'-OCH3; linkage at
      positions 34 and 35 is 3'-3'.
<400> 192
ugugaauauu agagagucug uauagcucua cccc
                                                                   34
<210> 193
<211> 34
<212> RNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<220>
<221> modified base
<222> (1)..(34)
<223> All pyrimidines are 2'F; a's and g's at positions
      2, 4-6, 11-16, 25 and 30 are 2'-OCH3; linkage at
      positions 34 and 35 is 3'-3'.
<400> 193
ugugaauauu agagagucug uauagcucua cccc
                                                                   34
<210> 194
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(42)
<223> A's and g's at positions 1-3 and 5 are 2'-OMe.
```

<220>

```
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 194
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                    42
<210> 195
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(42)
<223> A's and g's at positions 12 and 16-18 are 2'OMe.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 195
gggugccuuu ugccuagguu quqauuuqua accuucuqcc ca
                                                                    42
<210> 196
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(42)
<223> A's and g's at positions 21, 23-24 and 28 are
      2'-OMe.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 196
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                    42
<210> 197
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(42)
<223> A's and g's at positions 30-31, 38 and 42 are
      2'-OMe.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 197
```

```
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                    42
<210> 198
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(42)
<223> A's and g's at positions 1, 21, 23-24 and 28 are
      2'-OMe.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 198
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                    42
<210> 199
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(42)
<223> A's and g's at positions 2, 21, 23-24 and 28 are
      2'-OMe.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 199
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                    42
<210> 200
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(42)
<223> A's and g's at positions 3, 21, 23-24 and 28 are
      2'-OMe.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 200
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                    42
```

```
<210> 201
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(42)
<223> A's and g's at positions 5, 21, 23-24 and 28 are
      2'-OMe.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 201
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                    42
<210> 202
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(42)
<223> A's and g's at positions 12, 21, 23-24 and 28 are
      2'-OMe.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 202
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                    42
<210> 203
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(42)
<223> A's and g's at positions 16, 21, 23-24 and 28 are
      2'-OMe.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 203
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                    42
<210> 204
<211> 42
<212> RNA
```

```
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(42)
<223> A's and g's at positions 17, 21, 23-24 and 28 are
      2'-OMe.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 204
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                    42
<210> 205
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(42)
<223> A's and g's at positions 18, 21, 23-24, and 28 are
      2'-OMe.
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 205
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                    42
<210> 206
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<221> modified_base
<222> (1)..(42)
<223> A's and g's at positions 21, 23-24, 28 and 30 are
      2'-OMe.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                    42
<210> 207
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
```

```
<221> modified base
<222> (1)..(42)
<223> A's and g's at positions 21, 23-24, 28 and 31 are
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 207
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                   42
<210> 208
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(42)
<223> A's and g's at positions 21, 23-24, 28 and 38 are
      2'-OMe.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 208
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                   42
<210> 209
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(42)
<223> A's and g's at positions 21, 23-24, 28 and 42 are
      2'-OMe.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 209
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                   42
<210> 210
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(42)
<223> A's and g's at positions 1-3, 16-18, 21, 23-24,
```

```
28, 30-31 and 42 are 2'-OMe; linkage at positions
      42 and 43 is 3'-3'.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 210
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                    42
<210> 211
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<221> modified_base
<222> (1)..(42)
<223> A's and g's at positions 1-3, 16-18, 21, 23-24,
      28, 30 and 42 are 2'-OMe; linkage at positions 42
      and 43 is 3'-3'.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 211
gggugccuuu ugccuagguu gugauuugua accuucugcc ca
                                                                    42
<210> 212
<211> 41
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(41)
<223> A's and g's at positions 1-3, 16-18, 21, 23-24, 28
      and 30-31 are 2'-OMe; linkage at positions 41 and
      42 are 3'-3'.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 212
gggugccuuu ugccuagguu gugauuugua accuucugcc c
                                                                    41
<210> 213
<211> 37
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(37)
```

<223> A's and g's at positions 1-3, 16-18, 21, 23, 25-26

```
and 37 are 2'-OMe; linkage at positions 37 and 38
      is 3'-3'.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 213
gggugccuuu ugccuagguu guguaaccuu cugccca
                                                                    37
<210> 214
<211> 35
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(35)
<223> A's and g's at positions 1-3, 16-18, 21, 23-24 and
      35 are 2'-OMe; linkage at positions 35 and 36 is
      3'-3'.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 214
gggugccuuu ugccuagguu guaaccuucu gccca
                                                                    35
<210> 215
<211> 33
<212> RNA
<213> Artificial Sequence
<220>
<221> modified base
<222> (1)..(33)
<223> A's and g's at positions 1-3, 16-18, 21-22, and 33
      are 2'-OMe; linkage at positions 33 and 34 is
      3'-3'.
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
<400> 215
gggugccuuu ugccuagguu aaccuucugc cca
                                                                    33
<210> 216
<211> 42
<212> RNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Sequence
```

<400> 216
gggugccuuu ugccuagguu gugauuugua accuucugcc ca